1 1	CRF Errors Corrected by the STIC Systems Branch. CRF Processing Date: 5/20/200 Edited by: Verified by: Veri
	Changed a file from non-ASCII to ASCII
	Changed the margins in cases where the sequence text was wrapped flown to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of files page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	177.
	tal:
	Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop coden in artino acid sequences and adjusted the "(A)Length:" field accordingly (error
	Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in arrino acid sequences and adjusted the "(A)Length:" field accordingly (errodue to a Patentin bug). Sequences corrected:
	Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in arrino acid sequences and adjusted the "(A)Length:" field accordingly (errodue to a Patentin bug). Sequences corrected:



1600

RAW SEQUENCE LISTING DATE: 05/20/2003 PATENT APPLICATION: US/09/913,351A TIME: 08:36:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\1913351A.raw

3 <110> APPLICANT: DELGADO, AURORA BRIEVA VILLARRUBIA, VINCENTE GARCIA 5 GOMEZ-PAMO, ANTONIO GUERRERO RANIERI, JUAN PABLO PIVEL 7 GALLEGO, GUILLERMO GIMENEZ TUDURI, JOSE ANTONIO MATJI 10 <120> TITLE OF INVENTION: PHARMACOLOGICALLY ACTIVE POLYPEPTIDE GLYCOCONJUGATES 12 <130> FILE REFERENCE: 618999-1/JP/B-4275 14 <140> CURRENT APPLICATION NUMBER: 09/913,351A 15 <141> CURRENT FILING DATE: 1999-10-21 17 <150> PRIOR APPLICATION NUMBER: PCT/ES99/00338 18 <151> PRIOR FILING DATE: 1999-10-21 20 <150> PRIOR APPLICATION NUMBER: ES P9900408 21 <151> PRIOR FILING DATE: 1999-02-26 23 <160> NUMBER OF SEQ ID NOS: 5 25 <170> SOFTWARE: PatentIn Ver. 2.1 27 <210> SEQ ID NO: 1 28 <211> LENGTH: 230 29 <212> TYPE: PRT 30 <213> ORGANISM: Artificial Sequence 32 <220> FEATURE: 33 <223> OTHER INFORMATION: Description of Artificial Sequence: Formula sequence 36 <220> FEATURE: 37 <221> NAME/KEY: MOD RES 38 <222> LOCATION: (1)..(48) 39 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass 3-48 amino acids 42 <220> FEATURE: 43 <221> NAME/KEY: MOD_RES 44 <222> LOCATION: (50)..(62) 45 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass 9-13 amino acids 48 <220> FEATURE: 49 <221> NAME/KEY: MOD RES 50 <222> LOCATION: (64) 51 <223> OTHER INFORMATION: Gln, Glu, Arg or Lys 53 <220> FEATURE: 54 <221> NAME/KEY: MOD RES 55 <222> LOCATION: (65) 56 <223> OTHER INFORMATION: Variable amino acid 59 <220> FEATURE: 60 <221> NAME/KEY: MOD RES



DATE: 05/20/2003

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/913,351A TIME: 08:36:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\1913351A.raw

- 61 <222> LOCATION: (66)
- 62 <223> OTHER INFORMATION: Hydrophobic amino acid
- 64 <220> FEATURE:
- 65 <221> NAME/KEY: MOD RES
- 66 <222> LOCATION: (67)
- 67 <223> OTHER INFORMATION: Leu, Ile, Val or Met
- 69 <220> FEATURE:
- 70 <221> NAME/KEY: MOD RES
- 71 <222> LOCATION: (68)..(10,6)
- 72 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
- 73 15-39 amino acids
- 75 <220> FEATURE:
- 76 <221> NAME/KEY: MOD RES
- 77 <222> LOCATION: (109)
- 78 <223> OTHER INFORMATION: Hydrophilic amino acid
- 80 <220> FEATURE:
- 81 <221> NAME/KEY: MOD RES
- 82 <222> LOCATION: (110)
- 83 <223> OTHER INFORMATION: Gln, Glu or His
- 85 <220> FEATURE:
- 86 <221> NAME/KEY: MOD RES
- 87 <222> LOCATION: $(11\overline{1})$
- 88 <223> OTHER INFORMATION: Leu or Val
- 90 <220> FEATURE:
- 91 <221> NAME/KEY: MOD RES
- 92 <222> LOCATION: (112)..(117)
- 93 <223> OTHER INFORMATION: Variable amino acid
- 95 <220> FEATURE:
- 96 <221> NAME/KEY: MOD_RES
- 97 <222> LOCATION: (119)
- 98 <223> OTHER INFORMATION: Variable amino acid
- 100 <220> FEATURE:
- 101 <221> NAME/KEY: MOD_RES
- 102 <222> LOCATION: (121)..(122)
- 103 <223> OTHER INFORMATION: Variable amino acid
- 105 <220> FEATURE:
- 106 <221> NAME/KEY: MOD RES
- 107 <222> LOCATION: $(12\overline{3})$
- 108 <223> OTHER INFORMATION: Leu or Ile
- 110 <220> FEATURE:
- 111 <221> NAME/KEY: MOD RES
- 112 <222> LOCATION: $(12\overline{4})$...(179)
- 113 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
- 114 13-56 amino acids
- 117 <220> FEATURE:
- 118 <221> NAME/KEY: MOD RES
- 119 <222> LOCATION: (181)..(206)
- 120 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
- 121 15-26 amino acids



DATE: 05/20/2003 TIME: 08:36:51

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/913,351A

nout Cot . A.\ DEO AMC tot

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\I913351A.raw

```
123 <220> FEATURE:
 124 <221> NAME/KEY: MOD RES
 125 <222> LOCATION: (208)
 126 <223> OTHER INFORMATION: Variable amino acid
 128 <220> FEATURE:
 129 <221> NAME/KEY: MOD RES
 130 <222> LOCATION: (209)
 131 <223> OTHER INFORMATION: Val, Ile, Leu or Met
 133 <220> FEATURE:
 134 <221> NAME/KEY: MOD RES
 135 <222> LOCATION: (210)..(217)
 136 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
      1-8 amino acids
 137
 139 <220> FEATURE:
 140 <221> NAME/KEY: MOD RES
 141 <222> LOCATION: (218)..(230)
 142 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
    1-12 amino acids
 145 <400> SEQUENCE: 1
147 1
                       1.0
 159 65
              70
 164 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa
          100
                    105
 167 Xaa Xaa Xaa Xaa Cys Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                  120
 130
                135
                           140
 150
 165
                       170
 185
 195
                  200
 185 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
                215
```

188 Xaa Xaa Xaa Xaa Xaa

189 225



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/913,351A

DATE: 05/20/2003 TIME: 08:36:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\1913351A.raw

192 <210> SEO ID NO: 2 193 <211> LENGTH: 37 194 <212> TYPE: PRT 195 <213> ORGANISM: Ricinus communis 197 <400> SEQUENCE: 2 198 Glu Ser Lys Gly Glu Arg Glu Gly Ser Ser Ser Gln Gln Cys Arg Gln 201 Glu Val Gln Arg Lys Asp Leu Ser Ser Cys Glu Arg Tyr Leu Arg Gln 20 25 204 Ser Ser Ser Arg Arg 205 35 208 <210> SEQ ID NO: 3 209 <211> LENGTH: 68 210 <212> TYPE: PRT 211 <213> ORGANISM: Ricinus communis 213 <400> SEQUENCE: 3 214 Gln Gln Glu Ser Gln Gln Leu Gln Gln Cys Cys Asn Gln Val Lys 217 Gln Val Arg Asp Glu Cys Gln Cys Glu Ala Ile Lys Tyr Ile Ala Glu 20 25 220 Asp Gln Ile Gln Gln Gly Gln Leu His Gly Glu Glu Ser Glu Arg Val 40 223 Ala Gln Arg Ala Gly Glu Ile Val Ser Ser Cys Gly Val Arg Cys Met 226 Arg Gln Thr Arg 227 65 230 <210> SEQ ID NO: 4 231 <211> LENGTH: 34 232 <212> TYPE: PRT 233 <213> ORGANISM: Ricinus communis 235 <400> SEQUENCE: 4 236 Pro Ser Gln Gln Gly Cys Arg Gly Gln Ile Gln Glu Gln Gln Asn Leu 10 239 Arg Gln Cys Gln Glu Tyr Ile Lys Gln Gln Val Ser Gly Gln Gly Pro 240 242 Arg Arg 246 <210> SEQ ID NO: 5 247 <211> LENGTH: 65 248 <212> TYPE: PRT 249 <213> ORGANISM: Ricinus communis 251 <400> SEQUENCE: 5 252 Gln Glu Arg Ser Leu Arg Gly Cys Cys Asp His Leu Lys Gln Met Gln 255 Ser Gln Cys Arg Cys Glu Gly Leu Arg Gln Ala Ile Glu Gln Gln 258 Ser Gln Gly Gln Leu Gln Gly Gln Asp Val Phe Glu Ala Phe Arg Thr

261 Ala Ala Asn Leu Pro Ser Met Cys Gly Val Ser Pro Thr Glu Cys Arg

50

262

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/913,351A

DATE: 05/20/2003 TIME: 08:36:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\I913351A.raw

264 Phe 265 65



RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/913,351A

DATE: 05/20/2003 TIME: 08:36:52

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\I913351A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22-Seq#:1; Xaa Pos. 23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41 Seq#:1; Xaa Pos. 42,43,44,45,46,47,48,50,51,52,53,54,55,56,57,58,59,60,61-Seq#:1; Xaa Pos. 62,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79,80,81 Seq#:1; Xaa Pos. 82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99,100 Seq#:1; Xaa Pos. 101,102,103,104,105,106,109,110,111,112,113,114,115,116 Seq#:1; Xaa Pos. 117,119,121,122,123,124,125,126,127,128,129,130,131,132 Seq#:1; Xaa Pos. 133,134,135,136,137,138,139,140,141,142,143,144,145,146 Seq#:1; Xaa Pos. 147,148,149,150,151,152,153,154,155,156,157,158,159,160 Seq#:1; Xaa Pos. 161,162,163,164,165,166,167,168,169,170,171,172,173,174-Seq#:1; Xaa Pos. 175,176,177,178,179,181,182,183,184,185,186,187,188,189 Seq#:1; Xaa Pos. 190,191,192,193,194,195,196,197,198,199,200,201,202,203 Seq#:1; Xaa Pos. 204,205,206,208,209,210,211,212,213,214,215,216,217,219 Seq#:1; Xaa Pos. 220,221,222,223,224,225,226,227,228,229,230